

Figure 1/10

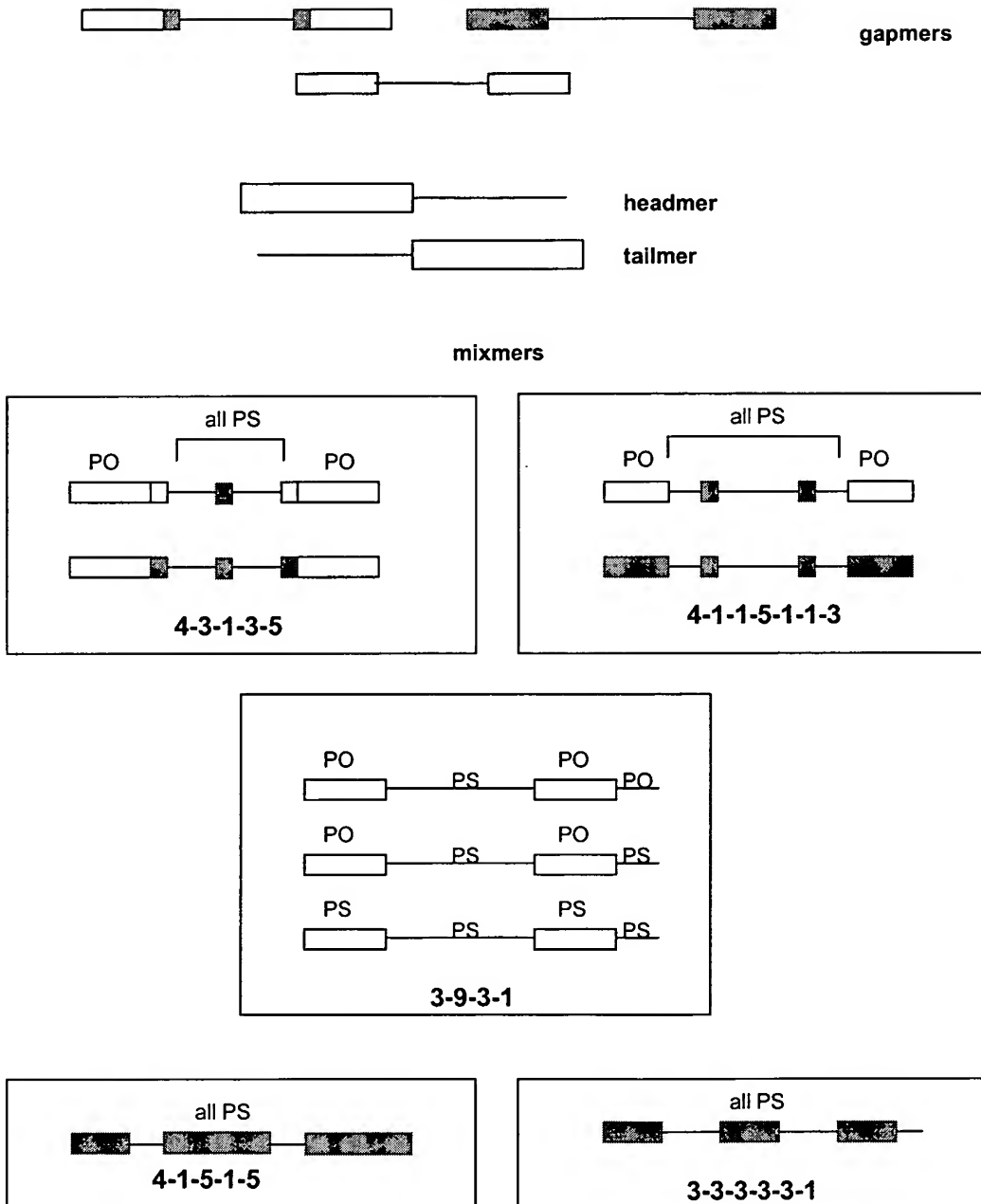
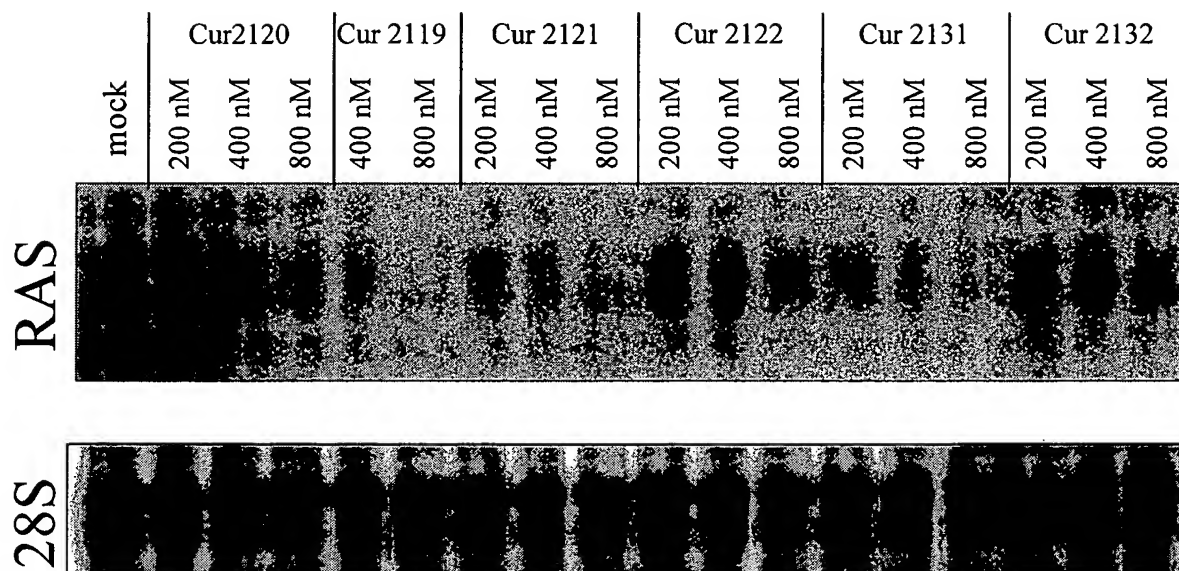


Figure 2/10



% mRNA relative to mock		200nM	400nM	800nM
Cur2119	PS20	-	35 (27)	21 (12)
Cur2120	PS20x6mm	80 (62)	83 (63)	53 (40)
Cur2121	LNA20	57 (43)	45 (30)	42 (26)
Cur2122	LNA20x6mm	88 (87)	65 (66)	52 (50)
Cur2131	LNA16	45 (44)	30 (26)	31 (16)
Cur2132	LNA16x6mm	145 (79)	119 (76)	110 (71)

Name	Composition	Sequence	Size	Relation to target
Cur 2119 (ISIS 2503)	PS-DNA	5'- tccgtcatcgctcctcaggg- 3'	20me r	Matched
Cur 2120 (ISIS control)	PS-DNA	5'- tcagtaatagccccacatgg-	20me r	6 mismatches

		3'		
Cur 2522	PS-DNA	5'-tccgtcatcgctcctc-3'	16mer	Matched
Cur 2131	PS-LNA gabmer	5'-TCCGtcacgcctCCTC-3'	16mer	Matched
Cur 2132	PS-LNA gabmer	5'-TCAGtaataagccCCAC-3'	16mer	5-mismatches

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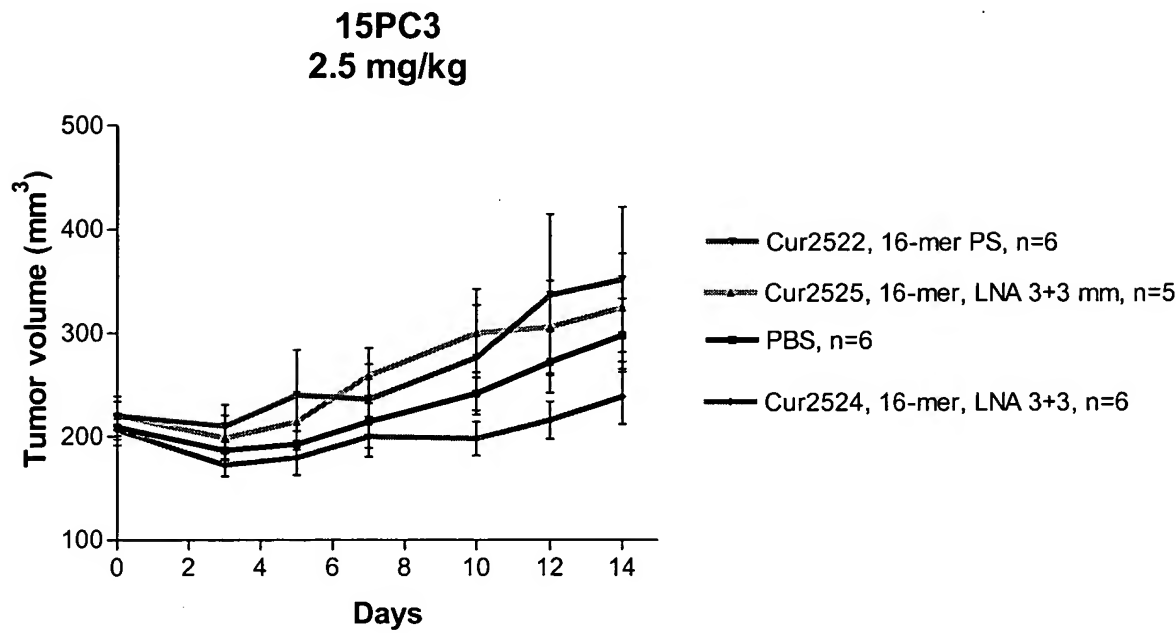


Figure 4/10

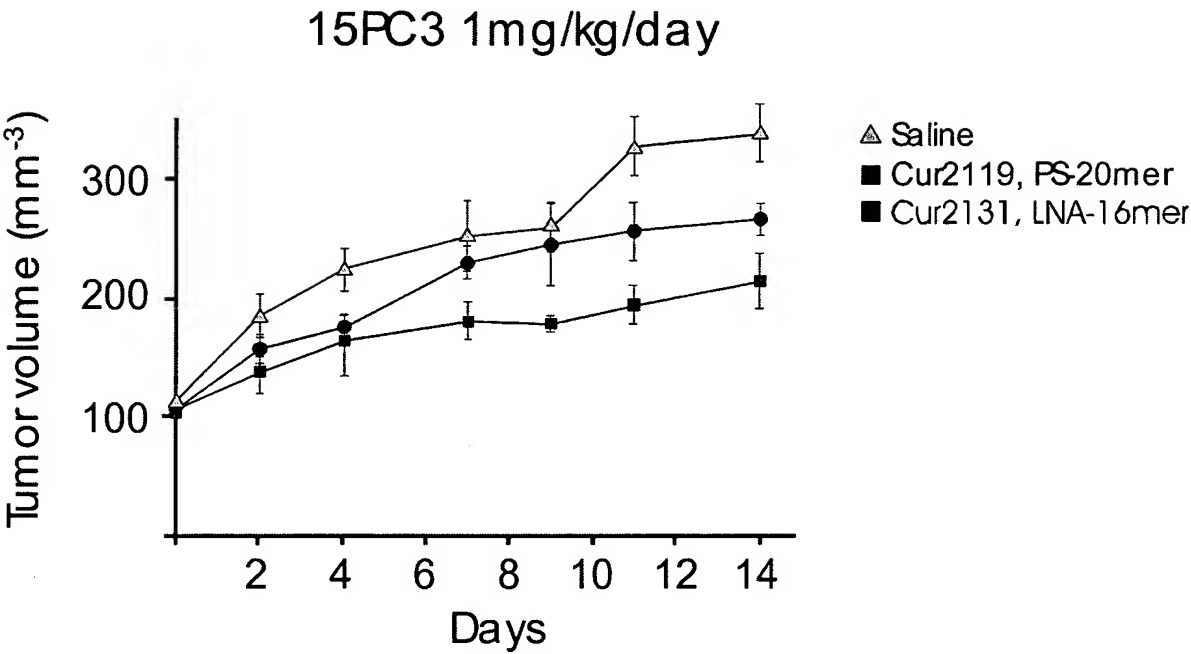
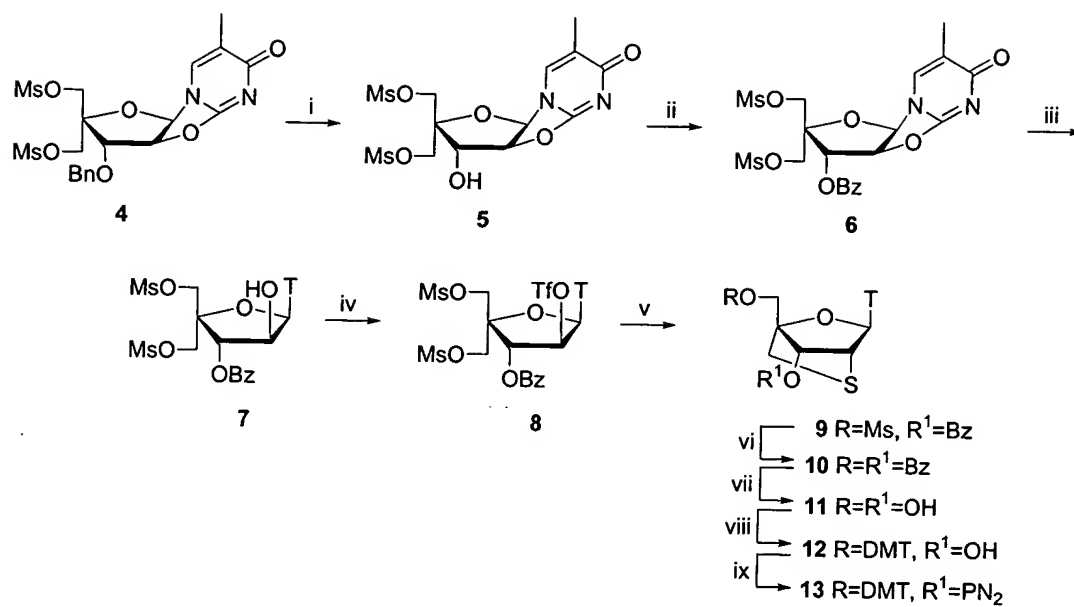
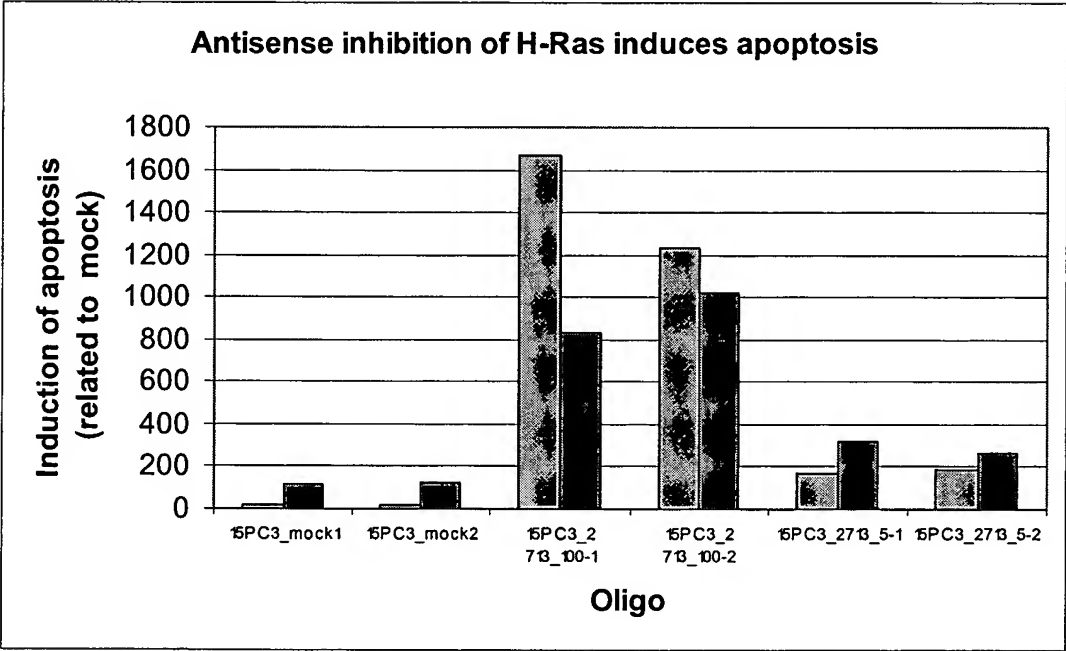


Figure 5/10



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Figure 6/10



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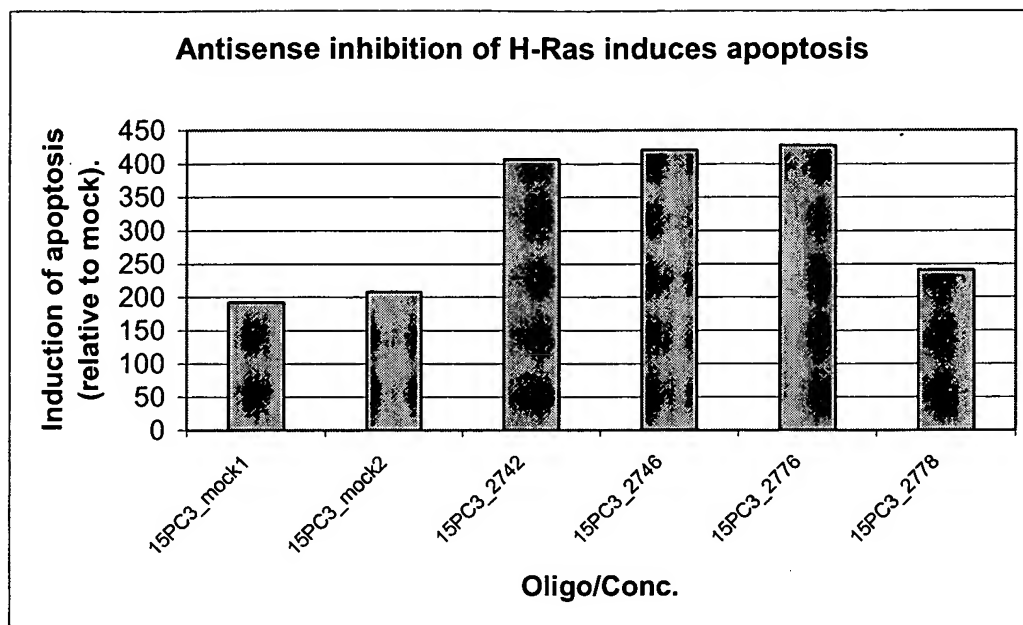


Figure 7/10

Seq ID NO: 1

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      6151   TTGAGGGAGG GTCTTGATGC CTGGGGTTAC CCGCAGAGGC CTGGGTGCCG
      AACTCCCTCC CAGAACTACG GACCCCAATG GCGGTCTCCG GACCCACGGC
15      6201   GGACGCTCCC CGGTTTGGCT GAAAGGAAAG CAGATGTGGT CAGCTTCTCC
      CCTGCGAGGG GCCAAACCGA CTTTCCTTTC GTCTACACCA GTCGAAGAGG
      6251   ACTGAGCCCA TCTGGTCTTC CCGGGGCTGG GCCCCATAGA TCTGGGTCCC
      TGA CTGCGGT AGACCAGAAG GGCCCCGACC CGGGGTATCT AGACCCAGGG
      6301   TGTGTGGCCC CCCTGGTCTG ATGCCGAGGA TACCCCTGCA AACTGCCAAT
20      ACACACCGGG GGGACCAGAC TACGGCTCCT ATGGGGACGT TTGACGGTTA
      6351   CCCAGAGGAC AAGACTGGGA AGTCCCTGCA GGGAGAGCCC ATCCCCGCAC
      GGGTCTCCTG TTCTGACCCT TCAGGGACGT CCCTCTCGGG TAGGGGCGTG
      6401   CCTGACCCAC AAGAGGGACT CCTGCTGCCC ACCAGGCATC CCTCCAGGGA
      GGACTGGGTG TTCTCCCTGA GGACGACGGG TGGTCCGTAG GGAGGTCCCT
25      6451   TCC
      AGG

```

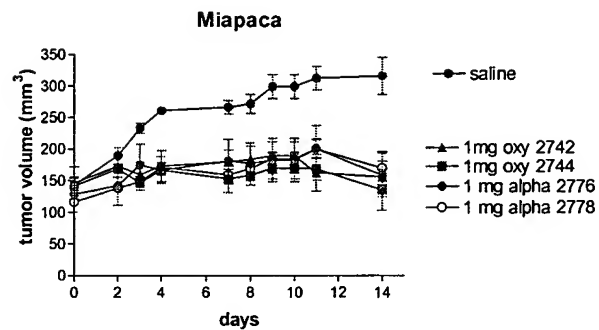
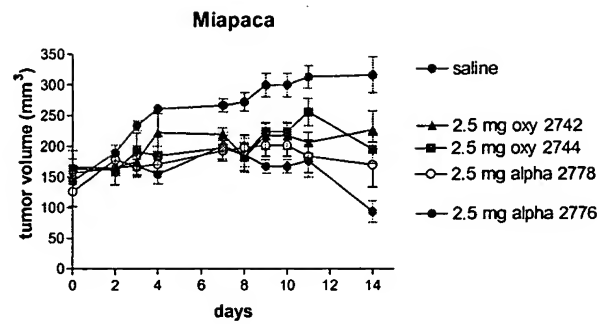
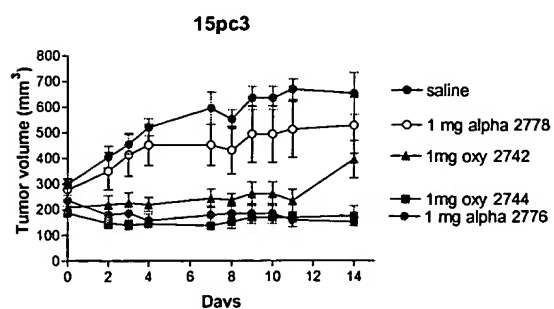
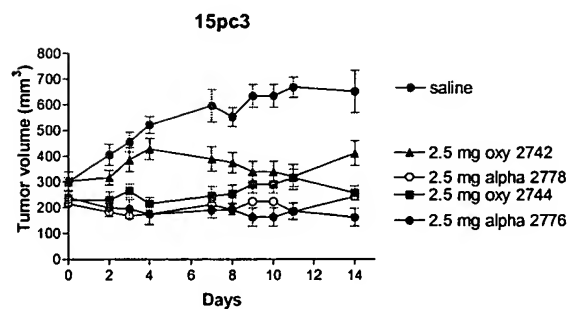


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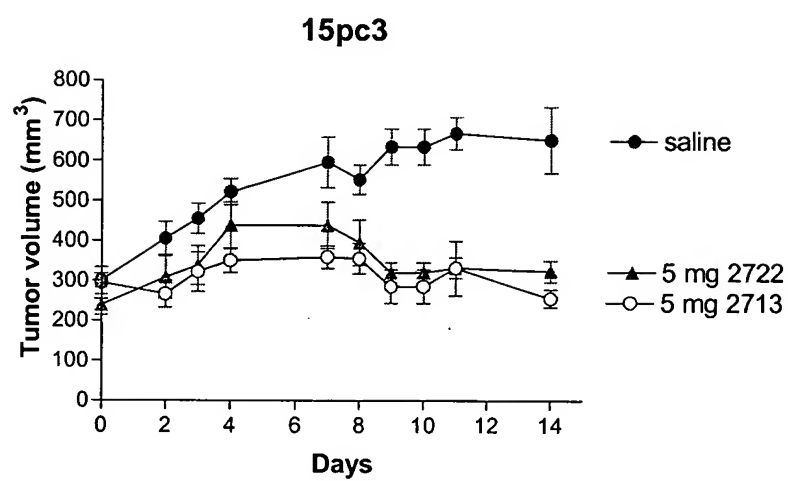
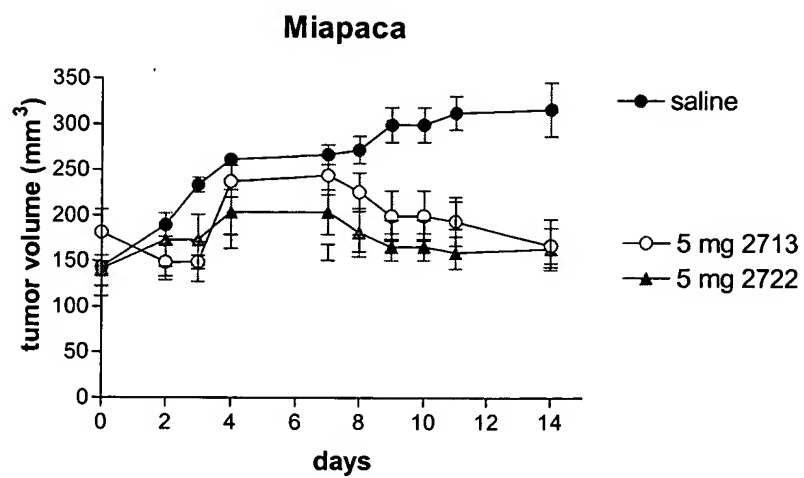
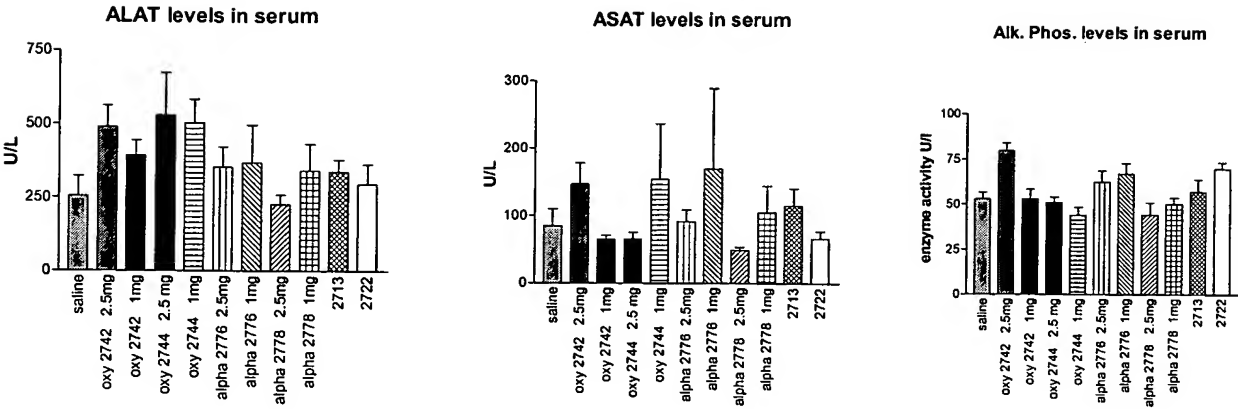


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